

1

1

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RESULT 4
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DEFINITION Sequence 15 from Patent WO0131003.
ACCESSION AX127740
VERSION AX127740.1 GI:14134387
KEYWORDS
SOURCE human.
ORGANISM Homo sapiens
 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
 Mammalia; Eutheria; Primates; Catartini; Homidae; Homo.
REFERENCE 1 (bases 1 to 2733)
AUTHORS Delneste, Y., Magistrelli, G., Jeannin, P. and Bonnefoy, J.Y.
TITLE Cloning, expression and characterisation of a gene expressed in
 tumour cells and involved in the regulation of the immune response
 Patent: WO 0131003-A 15 03-MAY-2001;
JOURNAL PIERRE FABRE MEDICAMENT (FR)
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BASE COUNT 656 a 783 c 687 g 607 t
 ORIGIN

Query Match 81.8%; Score 2726.6; DB 6; Length 2733;
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 Matches 2729; Conservative 0; Mismatches 4; Indels 0; Gaps 0;

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DEFINITION	Homo sapiens cDNA FLJ33085 f1s, clone TRACH2000420, weakly similar to MAJOR SURFACE-LABELLED TROPHOBLAST ANTIGEN PRECURSOR.				
ACCESSION	AK057647				

ACCESSION	AK057647
VERSION	AK057647.1
KEYWORDS	GI:16553406
SOURCE	oligo capping; fls (full insert sequence), Homo sapiens trachea cDNA to mRNA, clone_11b:TRACH2

ORGANISM

REFERENCE AUTHORS

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Katochka, R., Kikugawa, N., Kuroda, A., Sato, H., Kamata, R., Takami, S., Terashima, Y., Watanabe, M., Sugiyama, T., Irie, R., Otsuki, T., Sato, H., Ota, T., Wakamatsu, A., Inlil, S., Yamamoto, J., Isono, Y., Kawai-Hio, Y., Saito, R., Nishikawa, T., Kimura, K., Yamashita, H., Matsuo, K., Nakamura, Y., Sekine, M., Kikuchi, H., Kanda, K., Matsushima, M., Murakawa, K., Kanehori, K., Takahashi-Fujii, A., Oshima, A., Sugiyama, A., Kawakami, B., Suzuki, Y., Sugano, S., Nagashima, K., Mashino, Y., Nagai, K., and Isoyagi, T.

JOURNAL Unpublished
REFERENCE 2 (bases 1 to 2895)

AUTHORS Isogai, T., Otsuki, T. and Sugiyama, T.
TITLE Direct Submission
JOURNAL Submitted (24-OCT-2001) Takao Isogai

COMMENT

Economy, Trade and Industry of Japan; cDNA full insert sequencing: Research Association for Biotechnology (RAB); cDNA library construction: Helix Research Institute (HRI) (supported by Japan Key Technology Center etc.); 5' and 3' end one pass sequencing: RAB, HRI, and Biotechnology Center, National Institute of Technology and Evaluation; clone selection for full insert sequencing: RAB and HRI.

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ORIGIN

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1577	TGAACGTGAGCTCTACTTCAATGTTGGGTGTGAATTTAGAGCAACACTCCTGGGAGA	1636
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 ACCESSION AK026832.1
 VERSION AK026832.1
 KEYWORDS o1igo capping; fls (full insert sequence).
 SOURCE Homo sapiens human lung cDNA to mRNA, clone_l1b:LNG clone:LNG10890.
 ORGANISM Homo sapiens
 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Homidae; Homo.
 1 (sites)
 Kawakami, T., Noguchi, S., Itoh, T., Shigeta, K., Senba, T., Matsumura, K., Nakajima, Y., Mizuno, T., Morinaga, M., Tanigami, A., Fujiwara, T., Ono, T., Yamada, K., Fujii, Y., Ozaki, K., Hiro, M., Ohmori, Y., Ota, T., Suzuki, Y., Ohashi, M., Nishi, T., Shibahara, T., Tanaka, T., Nakamura, Y., Isogai, T. and Sugano, S.
 NEDO human cDNA sequencing project
 Unpublished
 2 (bases 1 to 2129)
 Sugano, S., Suzuki, Y., Ota, T., Ohashi, M., Nishi, T., Isogai, T., Shibahara, T., Tanaka, T. and Nakamura, Y.
 Direct Submission
 Submitted (29-AUG-2000) Sumio Sugano, Institute of Medical Science, University of Tokyo, Laboratory of Genome Structure Analysis, Human Genome Center, Shirokane-dai, 4-6-1, Minato-Ku, Tokyo 108-8639, Japan (E-mail:cdna@ims.u-tokyo.ac.jp, Tel:81-3-5449-5286, Fax:81-3-5449-5416)
 NEDO human cDNA sequencing project supported by Ministry of International Trade and Industry of Japan: cDNA full insert sequencing: Research Association for Biotechnology: cDNA library construction, 5'- & 3'-end one pass sequencing: Department of Virology and Human Genome Center, Institute of Medical Science, University of Tokyo (partly supported by Science and Technology Agency).
 FEATURES
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Matches 2109; Conservative 0; Mismatches 6; Indels 1; Gaps 1;

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RESULT 9
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 ACCESSION AB037745.1 GI:7243028
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 Mammalia; Eutheria; Primates; Catarrhini; Homiidae; Homo.
 REFERENCE 1 (sites)
 AUTHORS Nagase, T., Kikuno, R., Ishikawa, K.I., Hirosewa, M. and Ohara, O.
 TITLE Prediction of the coding sequences of unidentified human genes.
 XVI. The complete sequences of 150 new cDNA clones from brain which
 code for large proteins in vitro
 JOURNAL DNA Res. 7 (1), 65-73 (2000)
 MEDLINE 20181126
 REFERENCE 2 (bases 1 to 5567)
 AUTHORS Ohara, O., Nagase, T. and Kikuno, R.
 TITLE Direct Submission
 JOURNAL Submitted (31-JAN-2000) Osamu Ohara, Kazusa DNA Research Institute,
 Laboratory of DNA Technology, 1532-3 Yana, Kisarazu, Chiba
 292-0812, Japan (E-mail:cdna1to@kazusa.or.jp,
 URL: http://www.kazusa.or.jp/huge/, Tel:+81-438-52-3913,
 Fax:+81-438-52-3914)
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RESULT 10
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LOCUS AX013071
DEFINITION Sequence 11 from Patent W0954461.
ACCESSION AX013071
VERSION AX013071.1 GI:10040237
KEYWORDS
SOURCE
ORGANISM human.
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominiidae; Homo.
1 (bases 1 to 1717)
Schmitt,A., Specht,T., Dahl,E., Hinzmann,B., Rosenthal,A. and
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Human nucleic acid sequences of endometrium tumour tissue
Patent: WO 954461-A 11 28-OCT-1995;
SCHMITT ARMIN (DE); SPECHT THOMAS (DE); DAHL EDGAR (DE); HINZMANN
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Location/Qualifiers

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 VERSION AX127746.1 GI:14134393
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 SOURCE human.
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 REFERENCE
 1 (bases 1 to 1149)
 AUTHORS Delneste, Y., Magistrelli, G., Jeannin, P. and Bonnefoy, J. Y.
 TITLE Cloning, expression and characterization of a gene expressed in
 tumour cells and involved in the regulation of the immune response
 JOURNAL Patent: NO 0131003-A-21 03-MAY-2001;
 PIERRE FABRE MEDICAMENT (FR)
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Db      2157  TAACTCTACTTAATCTGAGAGAGAGTGTATTCAGTCCCAAGCAAGTCCCAAGCAGGATAC 2216
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Db      2217  CTGTGATGGGCTGTACCTTCATTCCTGTGGAGAGTGTCAAGCTGTGCCCTGTGTAC 2276
QY      2637  AGTGGCTGACTACCATGCTATCGTCAGCAGCTGTGTGGATCCGAGAGACTACTTA 2696
Db      2277  GGACCATGACTTCATGATGATGAGGAGCGCTGCAGAGAGATTTCAAGAACTTGTA 2336
QY      2697  CGTGGGAGAGAACCCAGCTATGCTGTGCGATTTCTGCTGTAGCAGAGAGTCAAC 2756
Db      2337  TGTGTGAATGAACCTTAATGTGCTATTAAGGAATTTCTTGCTGAGAAAAAGTTGCC 2396
QY      2757  CATCTGCAAAACCATAGATTTCTGGCTGAAGTGGGCATCTGTGAGGACCTGTACTGC 2816
Db      2397  AACCTGTGAACGGTGTGACTTTGGCTGAAGGTGGAGCCGGTGTGGAGCCTTTACTGC 2456
QY      2817  CATCTGCTCACCGTCTTGACTGTACTTTGGAAAAAGAAATCAAAACTAGAGTACAA 2876
Db      2457  CGTTTGTGCTGGTGGCTGTGACTGTCTGTGAAAAAGAAATCAAAACTGGAATACAA 2516
QY      2877  GTACTCCAAAGCTGTGATGAATGCTACTCTCAAGGACTGTGACCTGCCAGCAGTGAAC 2936
Db      2517  ATATTCCAAAGTATGATGAAGACTTAAGCAAGAGTGTGAACCTCCGGCTGCAGACAG 2576
QY      2937  CTGGCCCATCATGGAAGGAGAGATGTAGAGAGCAGCTCATCT- TTACCAAGCAAGAATC 2995
Db      2577  TTGTGCTATCATGGAAGAGAAATGAAGAGAGAGTGTATATTCATTAATTAACAGTTC 2636
QY      2996  ACTCTTTGGGAAGATCAATCATTTACTCAAGAGGACTCCTGATGATTTGACTCAGT 3055
Db      2637  ACTACTAGGAAAACTCAATCTTTGGCAACCAAGAAAAAGAGACCATTTTGAATCTGT 2696
QY      3056  GCCGCTGAAGACATC 3070
Db      2697  TCAACTGAAGAACTC 2711
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